

Figure 1

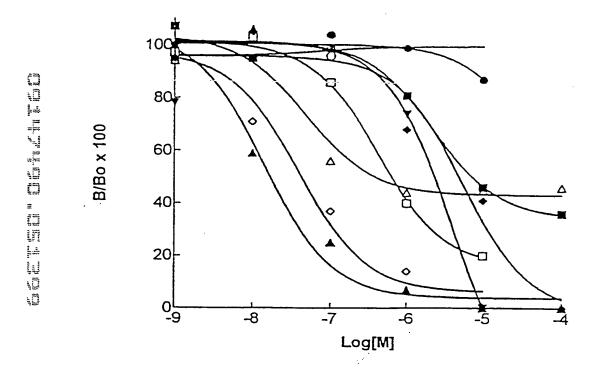


Figure 2



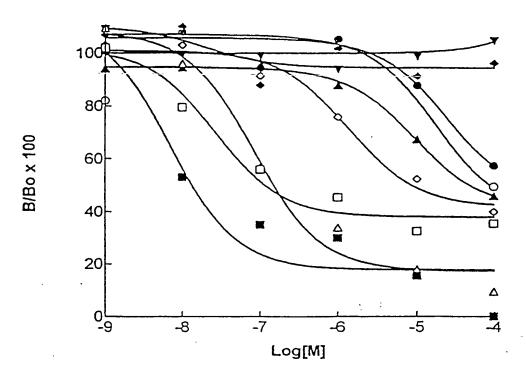


Figure 3

4/15

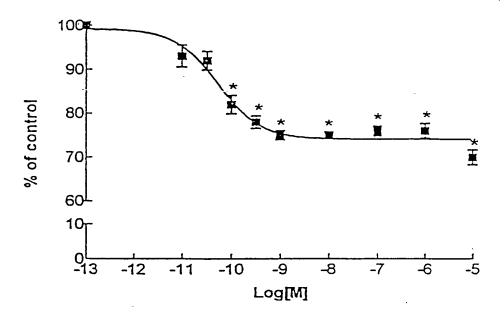


Figure 4

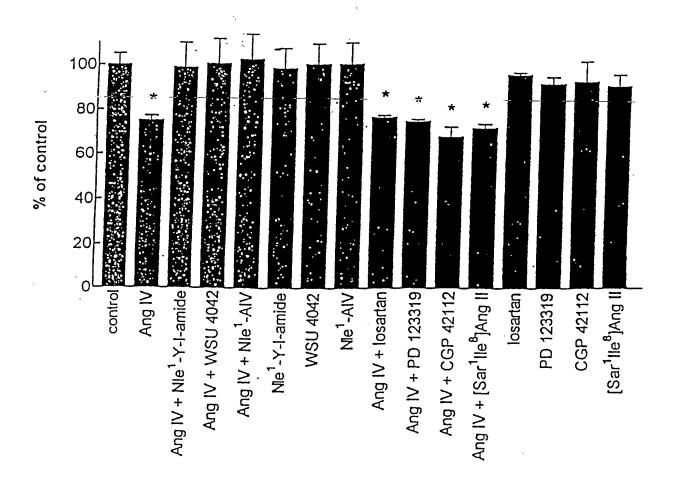


Figure 5

Figure 6

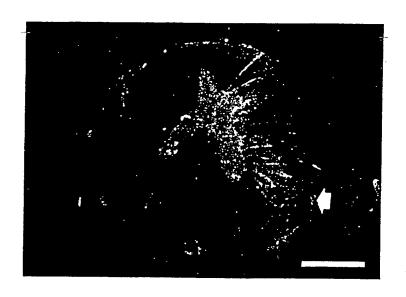


Figure 7

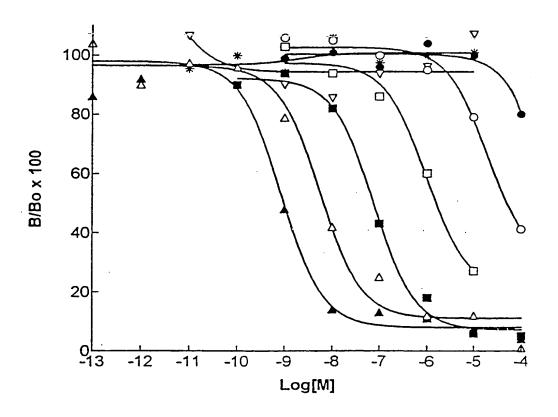


Figure 8

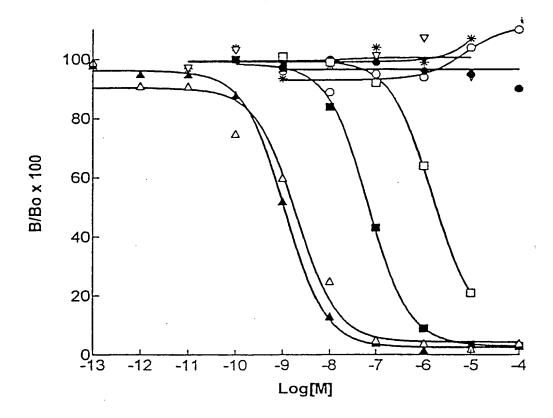


Figure 9

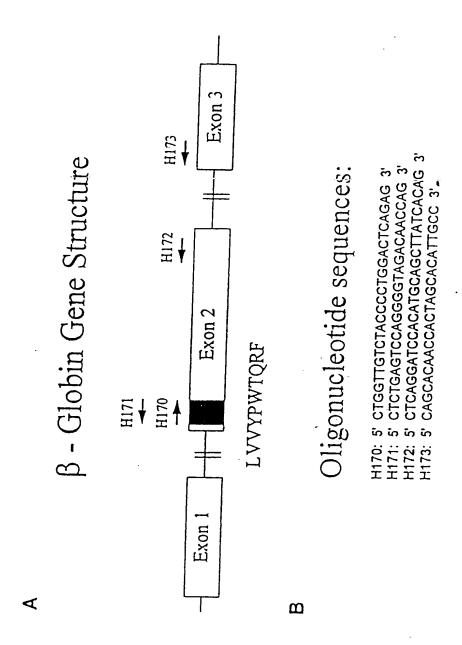


Figure 10

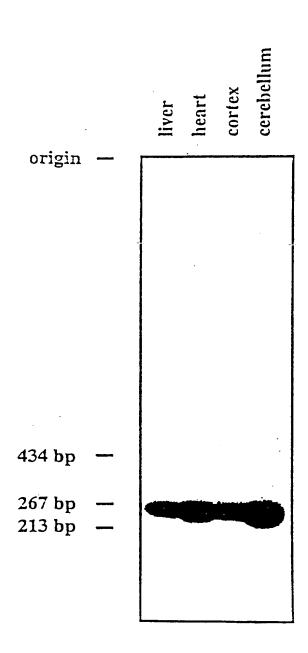


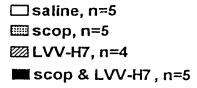
Figure 11

Figure 12

EΧ	10 20 30 CACAAACTCAGAAACAGACACCATGGTGCACCTGA M V H L
EX	40 50 60 70 80 90 CTGATGCTGAGAAGGCTGCTGTTAATGGCCTGTGGGGAAAGGTGAACCCTGATGATGTTG T D A E K A A V N G L W G K V N P D D V
EΧ	100 110 120 130 140 150 GTGGCGAGGCCTGGGCGGGCTGCTTGTCTCCCTTGGACCCAGAGGTACTTTGATA G G E A L G R L L V V Y P W T Q R Y F D
EX	160 170 180 190 200 210 GCTTTGGGGACCTGTCCTCTGCTATCATGGGTAACCCTAAGGTGAAGGCCCATG S F G D L S S A S A I M G N P K V K A H
ΕX	220 230 240 250 260 270 GCAAGAAGGTGATAAACGCCTCAATGATGGCCTGAAACACTTGGACAACCTCAAGGGCA G K K V I N A F N D G L K H L D N L K G
EX	280 290 300 310 320 330 CCTTTGCTCATCTGAGTGAACTCCACTGTGACAAGCTGCATGTGGATCCTGAGAACTTCA T F A H L S E L H C D K L H V D P E N F
EX	340 350 360 370 380 390 GGCTCCTGGGCAATATGATTGTGTTGTGTGGGCCACCACCTGGGCAAGGAATTCACCC R L L G N M I V I V L G H H L G K E F T
EX	400 410 420 430 440 450 CCTGTGCACAGGCTGCCTCACA
EX	P C A Q A A F Q K V V A G V A S A L A H 460 470 480 490 500 510 AGTACCACTÁAACCTCTTTTCCTGCTCTTTTTTTTTTTTT
EX	520 530 540 550 560 570 GAGCATCTGTCAGTTGTCAAAATGACAAAGACCTTTGAAAATCTGTCCTACTAATAA
EΧ	580 590 600 610 AAGGCATTTACTTTCACTGCAAAAAAAAAAAAAAAAAA

Figure 13

14/15



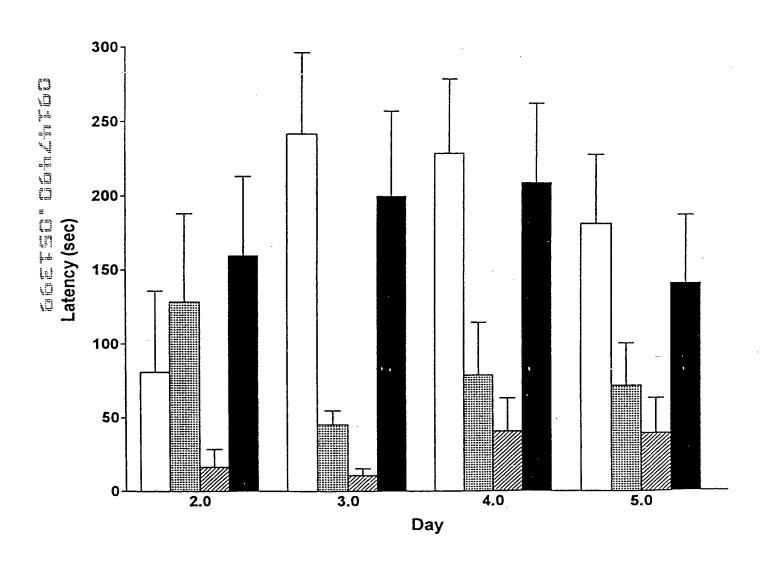
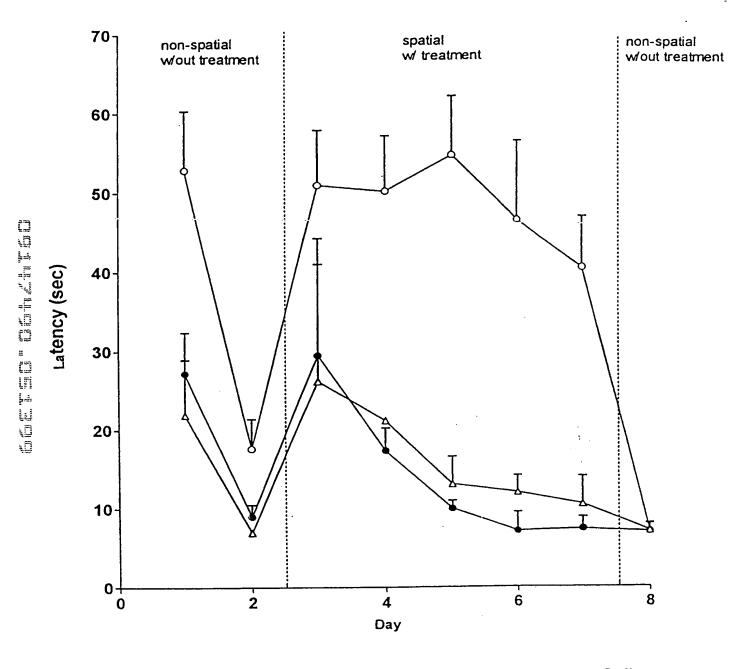


Figure 14

15/15



-<u></u>--Saline

-->-Scop

-Scop & LVV-H7

Figure 15